RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/524,198_
Source: _PCT/10_
Date Processed by STIC: _2/26/05_

ENTERED

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        RAW SEQUENCE LISTING                    DATE: 02/26/2005
        PATENT APPLICATION:  US/10/524,198      TIME: 10:44:18

        Input Set : A:\2002.013 SEQ LIST.txt
        Output Set: N:\CRF4\02262005\J524198.raw


      3 <110> APPLICANT: AKZO Nobel N.V.
      5 <120> TITLE OF INVENTION: Novel mastitis vaccine
      7 <130> FILE REFERENCE: 2002.013
C-->  9 <140> CURRENT APPLICATION NUMBER: US/10/524,198
C-->  9 <141> CURRENT FILING DATE: 2005-02-10
      9 <160> NUMBER OF SEQ ID NOS: 2
     11 <170> SOFTWARE: PatentIn version 3.1
     13 <210> SEQ ID NO: 1
     14 <211> LENGTH: 603
     15 <212> TYPE: DNA
     16 <213> ORGANISM: Streptococcus uberis
     18 <220> FEATURE:
     19 <221> NAME/KEY: CDS
     20 <222> LOCATION: (1)..(600)
     21 <223> OTHER INFORMATION:
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     25 atg ttt aaa ttt tta aag cgt gtt gtt ttt cta gct ttt ctg att ttt      48
     26 Met Phe Lys Phe Leu Lys Arg Val Val Phe Leu Ala Phe Leu Ile Phe
     27 1               5                   10                  15
     29 tgt ttt tat caa gct tat ata aca cat caa aat gta caa aat gtc atg      96
     30 Cys Phe Tyr Gln Ala Tyr Ile Thr His Gln Asn Val Gln Asn Val Met
     31             20                  25                  30
     33 caa tac aaa cca atg gtt gaa aaa acc ttg gct gaa aat gat acg act     144
     34 Gln Tyr Lys Pro Met Val Glu Lys Thr Leu Ala Glu Asn Asp Thr Thr
     35             35                  40                  45
     37 gcc aat gtc aat tta gtt tta gca atg atc tac aca gaa aca aaa ggt     192
     38 Ala Asn Val Asn Leu Val Leu Ala Met Ile Tyr Thr Glu Thr Lys Gly
     39     50                  55                  60
     41 ggt cag gca gat gtc atg caa tct agc gaa agt agt agt ggt gtg act     240
     42 Gly Gln Ala Asp Val Met Gln Ser Ser Glu Ser Ser Ser Gly Val Thr
     43 65                  70                  75                  80
     45 aac tca att acc gac agt caa tct agt att caa cac ggt gtc aaa ctc     288
     46 Asn Ser Ile Thr Asp Ser Gln Ser Ser Ile Gln His Gly Val Lys Leu
     47             85                  90                  95
     49 ttg tct gag aat ttg act tta gct gag aaa gct gga gta gac tct tgg     336
     50 Leu Ser Glu Asn Leu Thr Leu Ala Glu Lys Ala Gly Val Asp Ser Trp
     51             100                 105                 110
     53 act gca gta caa gct tac aat ttt gga aca gct tac att gat tat gtg     384
     54 Thr Ala Val Gln Ala Tyr Asn Phe Gly Thr Ala Tyr Ile Asp Tyr Val
     55             115                 120                 125
     57 gca aaa aat ggt ggt gac aac act atc tct ttg gct agt cat tat tct     432
     58 Ala Lys Asn Gly Gly Asp Asn Thr Ile Ser Leu Ala Ser His Tyr Ser
     59     130                 135                 140
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61 aaa agt gtt gta gct cca agt tta ggg aat aag gat gga aaa atg tat        480
62 Lys Ser Val Val Ala Pro Ser Leu Gly Asn Lys Asp Gly Lys Met Tyr
63 145                 150                 155                 160
65 tta tat tac cat cca att gcc ctc ctc tat ggc ggt aaa ctt tat caa        528
66 Leu Tyr Tyr His Pro Ile Ala Leu Leu Tyr Gly Gly Lys Leu Tyr Gln
67                 165                 170                 175
69 aat ggt ggt aat att tat tat tca cga gaa gtt cat ttt aat tat tac .      576
70 Asn Gly Gly Asn Ile Tyr Tyr Ser Arg Glu Val His Phe Asn Tyr Tyr
71                 180                 185                 190
73 ctc ata caa tta tta tct aaa ttt taa                                    603
74 Leu Ile Gln Leu Leu Ser Lys Phe
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78 <210> SEQ ID NO: 2
79 <211> LENGTH: 200
80 <212> TYPE: PRT
81 <213> ORGANISM: Streptococcus uberis

83 <400> SEQUENCE: 2

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86 1               5                   10                  15
89 Cys Phe Tyr Gln Ala Tyr Ile Thr His Gln Asn Val Gln Asn Val Met
90              20                  25                  30
93 Gln Tyr Lys Pro Met Val Glu Lys Thr Leu Ala Glu Asn Asp Thr Thr
94          35                  40                  45
97 Ala Asn Val Asn Leu Val Leu Ala Met Ile Tyr Thr Glu Thr Lys Gly
98      50                  55                  60
101 Gly Gln Ala Asp Val Met Gln Ser Ser Glu Ser Ser Ser Gly Val Thr
102 65                  70                  75                  80
105 Asn Ser Ile Thr Asp Ser Gln Ser Ser Ile Gln His Gly Val Lys Leu
106                 85                  90                  95
109 Leu Ser Glu Asn Leu Thr Leu Ala Glu Lys Ala Gly Val Asp Ser Trp
110                 100                 105                 110
113 Thr Ala Val Gln Ala Tyr Asn Phe Gly Thr Ala Tyr Ile Asp Tyr Val
114         115                 120                 125
117 Ala Lys Asn Gly Gly Asp Asn Thr Ile Ser Leu Ala Ser His Tyr Ser
118     130                 135                 140
121 Lys Ser Val Val Ala Pro Ser Leu Gly Asn Lys Asp Gly Lys Met Tyr
122 145                 150                 155                 160
125 Leu Tyr Tyr His Pro Ile Ala Leu Leu Tyr Gly Gly Lys Leu Tyr Gln
126                 165                 170                 175
129 Asn Gly Gly Asn Ile Tyr Tyr Ser Arg Glu Val His Phe Asn Tyr Tyr
130                 180                 185                 190
133 Leu Ile Gln Leu Leu Ser Lys Phe
134         195                 200
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VERIFICATION SUMMARY   DATE: 02/26/2005
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:24 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:21